

## SEQUENCE LISTING

<110> Jerome B. Posner  
 Josep O. Dalmau  
 Myrna R. Rosenfeld

<120> Ma FAMILY POLYPEPTIDES AND ANTI-Ma  
 ANTIBODIES

<130> 2581.1004-004

<150> 09/189,527

<151> 1998-11-10

<160> 14

<170> FastSEQ for Windows Version 4.0

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<211> 18

<212> DNA

<213> homo sapiens

<400> 1

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18

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<211> 19

<212> DNA

<213> homo sapiens

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gtctttgcgg atgtccacg

19

<210> 3

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<213> homo sapiens

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<221> misc\_feature

<222> 1699

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 cagcgtcccc gagctcggct ccgagtgcac ctacggactg actgtggggg cagagaaggg 120  
 cgagatcagg actctgtcct tgtaatcgt gactgcatga aggtcgcctc cctcgggcct 180  
 acttggtggg agtgtctggt attgttctaa ggccaggagc acggtgagcc acagtctgtt 240

204070-09846001

|   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
|---|--|--|--|--|--|--|--|--|--|--|--|--|--|--|-----|
| ggtagaattt ggcgtcttga tagttgagaa a atg gcg atg aca ctg ttg gaa  |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 292 |
| Met Ala Met Thr Leu Leu Glu                                     |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 1 5   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| gac tgg tgc cgg ggg atg gat gtg aac tcc cag aga act ctg tta gtc |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 340 |
| Asp Trp Cys Arg Gly Met Asp Val Asn Ser Gln Arg Thr Leu Leu Val |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 10 15 20  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| tgg ggc atc cca gtg aac tgt gat gag gct gaa atc gaa gag acc ctc |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 388 |
| Trp Gly Ile Pro Val Asn Cys Asp Glu Ala Glu Ile Glu Glu Thr Leu |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 25 30 35  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| cag gct gcg atg ccc cag gtc tcc tac cga atg ctt ggg aga atg ttc |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 436 |
| Gln Ala Ala Met Pro Gln Val Ser Tyr Arg Met Leu Gly Arg Met Phe |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 40 45 50 55   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| tgg agg gaa gaa aat gcg aaa gca gcc tta tta gag ctc act ggc gct |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 484 |
| Trp Arg Glu Glu Asn Ala Lys Ala Ala Leu Leu Glu Leu Thr Gly Ala |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 60 65 70  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| gta gat tac gcc gcg atc ccc agg gag atg ccg ggc aaa gga ggg gtc |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 532 |
| Val Asp Tyr Ala Ala Ile Pro Arg Glu Met Pro Gly Lys Gly Gly Val |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 75 80 85  |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| tgg aaa gtg tta ttt aag ccc cca act tct gat gct gaa ttt tta gaa |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 580 |
| Trp Lys Val Leu Phe Lys Pro Pro Thr Ser Asp Ala Glu Phe Leu Glu |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 90 95 100   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| aga ttg cac ctc ttc cta gct aga gag ggg tgg acc gtg caa gat gtt |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 628 |
| Arg Leu His Leu Phe Leu Ala Arg Glu Gly Trp Thr Val Gln Asp Val |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 105 110 115   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| gcc cgt gtc ctt ggg ttt cag aac cct act ccg acc ccg ggc cca gag |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 676 |
| Ala Arg Val Leu Gly Phe Gln Asn Pro Thr Pro Thr Pro Gly Pro Glu |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 120 125 130 135   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| atg cca gca gag atg cta aac tat att ttg gat aat gtt att cag cct |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 724 |
| Met Pro Ala Glu Met Leu Asn Tyr Ile Leu Asp Asn Val Ile Gln Pro |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 140 145 150   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| ctt gtt gag tcc ata tgg tac aag agg ctg aca ctt ttc tcg ggg aag |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 772 |
| Leu Val Glu Ser Ile Trp Tyr Lys Arg Leu Thr Leu Phe Ser Gly Lys |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 155 160 165   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| gga cat ccc agg gcc tgg aga gga aac ttt gat ccc tgg ctg gag cac |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 820 |
| Gly His Pro Arg Ala Trp Arg Gly Asn Phe Asp Pro Trp Leu Glu His |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 170 175 180   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| act aat gag gtc cta gag gag tgg cag gtg tcc gat gta gaa aag agg |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 868 |
| Thr Asn Glu Val Leu Glu Glu Trp Gln Val Ser Asp Val Glu Lys Arg |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 185 190 195   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| cgg cgg ttg atg gag agt ctt aga ggc ccc gcc gct gat gtt att cgc |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 916 |
| Arg Arg Leu Met Glu Ser Leu Arg Gly Pro Ala Ala Asp Val Ile Arg |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 200 205 210 215   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| atc ctt aag tcc aac aac ccc gcg ata acc act gcc gaa tgc ctg aag |  |  |  |  |  |  |  |  |  |  |  |  |  |  | 964 |
| Ile Leu Lys Ser Asn Asn Pro Ala Ile Thr Thr Ala Glu Cys Leu Lys |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |
| 220 225 230   |  |  |  |  |  |  |  |  |  |  |  |  |  |  |     |

gcg ctt gag cag gtg ttt ggg agc gtt gag agc tct agg gat gcc cag 1012  
 Ala Leu Glu Gln Val Phe Gly Ser Val Glu Ser Ser Arg Asp Ala Gln  
                   235                  240                  245

atc aaa ttt ctg aac act tat cag aac ccg gga gaa aaa ttg tct gct 1060  
 Ile Lys Phe Leu Asn Thr Tyr Gln Asn Pro Gly Glu Lys Leu Ser Ala  
                   250                  255                  260

tat gtc att cgt ctg gag cct ctg cta cag aag gtg gta gag aag ggg 1108  
 Tyr Val Ile Arg Leu Glu Pro Leu Leu Gln Lys Val Val Glu Lys Gly  
                   265                  270                  275

gcc att gat aaa gat aat gtg aac cag gcc cgc cta gag cag gtc att 1156  
 Ala Ile Asp Lys Asp Asn Val Asn Gln Ala Arg Leu Glu Gln Val Ile  
                   280                  285                  290                  295

gcc ggg gcc aac cac agc ggg gcc atc cga agg cag ctg tgg ctt acc 1204  
 Ala Gly Ala Asn His Ser Gly Ala Ile Arg Arg Gln Leu Trp Leu Thr  
                   300                  305                  310

ggg gct ggg gaa ggg cca ggc ccc aaa cct ctt tca gtt gct ggt gca 1252  
 Gly Ala Gly Glu Gly Pro Gly Pro Lys Pro Leu Ser Val Ala Gly Ala  
                   315                  320                  325

gat ccg tgaggaggaa gcccagggag gaggaggagg aggctgaggc cacccttctg 1308  
 Asp Pro

cagttaggcc tggaagggca cttctgagtg ccaggaaagg cagctttagt gcagacctag 1368  
 atcacagcta cttttcttgt ccctgtgggg tcttacagat gtgtctctga gtagtaaagg 1428  
 cttagccttg ttctgttttg ttgttttttg gaggggaagg ttagtcaggc ctgagtattc 1488  
 atgtaacatt ctaaaattgt gccagcgagc accgtgaacg actgcaatgc aagcgggtct 1548  
 tgctggctaa aatgcccagg taaaggggtg gttggacaca gcgcttagtg cacgctgtca 1608  
 tcatggacat cataatcagt tgtgaaaaac acgcgaacct atgacacttc ttattccaca 1668  
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 cagtaaaagt atgcactgtt agattactga taacgcggat agatttttgt ttaccataaa 1788  
 ttgttccaga tttatatata tggaaggaag tgtgcattta ttagctatta ctcaacttta 1848  
 caatgcaaac atcttatttc tcatctttta acatgtcgac cagtttaatt gaaaagtatt 1908  
 ctgagactgc aaaatggggt gttaaaaaat actgcagtta cggagctgtg taaaccagtt 1968  
 tctcattgca taagatacag atgtaaattg catggagagg ttgatatgca cctgtacagt 2028  
 aattcactcc cccatttcac ttctttgtca gagaatagtt cttgttcata ctgagtgttc 2088  
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<210> 4

<211> 329

<212> PRT

<213> homo sapiens

<400> 4

Met Ala Met Thr Leu Leu Glu Asp Trp Cys Arg Gly Met Asp Val Asn  
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 Ser Gln Arg Thr Leu Leu Val Trp Gly Ile Pro Val Asn Cys Asp Glu  
                   20                  25                  30  
 Ala Glu Ile Glu Glu Thr Leu Gln Ala Ala Met Pro Gln Val Ser Tyr  
                   35                  40                  45  
 Arg Met Leu Gly Arg Met Phe Trp Arg Glu Glu Asn Ala Lys Ala Ala  
                   50                  55                  60

1003850-010400

Leu Leu Glu Leu Thr Gly Ala Val Asp Tyr Ala Ala Ile Pro Arg Glu  
 65 70 75 80  
 Met Pro Gly Lys Gly Gly Val Trp Lys Val Leu Phe Lys Pro Pro Thr  
 85 90 95  
 Ser Asp Ala Glu Phe Leu Glu Arg Leu His Leu Phe Leu Ala Arg Glu  
 100 105 110  
 Gly Trp Thr Val Gln Asp Val Ala Arg Val Leu Gly Phe Gln Asn Pro  
 115 120 125  
 Thr Pro Thr Pro Gly Pro Glu Met Pro Ala Glu Met Leu Asn Tyr Ile  
 130 135 140  
 Leu Asp Asn Val Ile Gln Pro Leu Val Glu Ser Ile Trp Tyr Lys Arg  
 145 150 155 160  
 Leu Thr Leu Phe Ser Gly Lys Gly His Pro Arg Ala Trp Arg Gly Asn  
 165 170 175  
 Phe Asp Pro Trp Leu Glu His Thr Asn Glu Val Leu Glu Glu Trp Gln  
 180 185 190  
 Val Ser Asp Val Glu Lys Arg Arg Arg Leu Met Glu Ser Leu Arg Gly  
 195 200 205  
 Pro Ala Ala Asp Val Ile Arg Ile Leu Lys Ser Asn Asn Pro Ala Ile  
 210 215 220  
 Thr Thr Ala Glu Cys Leu Lys Ala Leu Glu Gln Val Phe Gly Ser Val  
 225 230 235 240  
 Glu Ser Ser Arg Asp Ala Gln Ile Lys Phe Leu Asn Thr Tyr Gln Asn  
 245 250 255  
 Pro Gly Glu Lys Leu Ser Ala Tyr Val Ile Arg Leu Glu Pro Leu Leu  
 260 265 270  
 Gln Lys Val Val Glu Lys Gly Ala Ile Asp Lys Asp Asn Val Asn Gln  
 275 280 285  
 Ala Arg Leu Glu Gln Val Ile Ala Gly Ala Asn His Ser Gly Ala Ile  
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 Arg Arg Gln Leu Trp Leu Thr Gly Ala Gly Glu Gly Pro Gly Pro Lys  
 305 310 315 320  
 Pro Leu Ser Val Ala Gly Ala Asp Pro  
 325

<210> 5  
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<400> 5  
 gggaatggcc gagacatc

18

<210> 6  
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<220>  
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 <222> (1)...(585)

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 Pro Leu Ala Leu Leu Glu Asp Trp Cys Arg Ile Met Ser Val Asp Glu  
 1 5 10 15

cag aag tca ctg atg gtt acg ggg ata ccg gcg gac ttt gag gag gct 96

Gln Lys Ser Leu Met Val Thr Gly Ile Pro Ala Asp Phe Glu Glu Ala  
                   20                                  25                                  30

gag att cag gag gtc ctt cag gag act tta aag tct ctg ggc agg tat 144  
 Glu Ile Gln Glu Val Leu Gln Glu Thr Leu Lys Ser Leu Gly Arg Tyr  
                   35                                  40                                  45

aga ctg ctt ggc aag ata ttc cgg aag cag gag aat gcc aat gct gtc 192  
 Arg Leu Leu Gly Lys Ile Phe Arg Lys Gln Glu Asn Ala Asn Ala Val  
                   50                                  55                                  60

tta cta gag ctt ctg gaa gat act gat gtc tcg gcc att ccc agt gag 240  
 Leu Leu Glu Leu Leu Glu Asp Thr Asp Val Ser Ala Ile Pro Ser Glu  
                   65                                  70                                  75                                  80

gtc cag gga aag ggg ggt gtc tgg aaa gtg atc ttt aag acc cct aat 288  
 Val Gln Gly Lys Gly Gly Val Trp Lys Val Ile Phe Lys Thr Pro Asn  
                                   85                                  90                                  95

cag gac act gag ttt ctt gaa aga ttg aac ctg ttt cta gaa aaa gag 336  
 Gln Asp Thr Glu Phe Leu Glu Arg Leu Asn Leu Phe Leu Glu Lys Glu  
                                   100                                  105                                  110

ggg cag acg gtc tcg ggt atg ttt cga gcc ctg ggg cag gag gcg ttg 384  
 Gly Gln Thr Val Ser Gly Met Phe Arg Ala Leu Gly Gln Glu Ala Leu  
                                   115                                  120                                  125

tct cca gcc aca gtg ccc tgc atc tca cca gaa tta ctg gcc cat ttg 432  
 Ser Pro Ala Thr Val Pro Cys Ile Ser Pro Glu Leu Leu Ala His Leu  
                                   130                                  135                                  140

ttg gga cag gca atg gca cat gcg cct cag ccc ctg cta ccc atg aga 480  
 Leu Gly Gln Ala Met Ala His Ala Pro Gln Pro Leu Leu Pro Met Arg  
                                   145                                  150                                  155                                  160

tac cgg aaa ctg cga gta ttc tca ggg agt gct gtc cca gcc cca gag 528  
 Tyr Arg Lys Leu Arg Val Phe Ser Gly Ser Ala Val Pro Ala Pro Glu  
                                   165                                  170                                  175

gaa gag tcc ttt gag gtc tgg ttg gaa cag gcc acg gag ata gtc aaa 576  
 Glu Glu Ser Phe Glu Val Trp Leu Glu Gln Ala Thr Glu Ile Val Lys  
                                   180                                  185                                  190

gag tgg cct tgaacacaac caaaaaaaaaa aaaaaaaaaag 615  
 Glu Trp Pro  
                   195

&lt;210&gt; 7

&lt;211&gt; 195

&lt;212&gt; PRT

&lt;213&gt; homo sapiens

&lt;400&gt; 7

Pro Leu Ala Leu Leu Glu Asp Trp Cys Arg Ile Met Ser Val Asp Glu  
   1                                  5                                  10                                  15  
 Gln Lys Ser Leu Met Val Thr Gly Ile Pro Ala Asp Phe Glu Glu Ala  
                   20                                  25                                  30

Glu Ile Gln Glu Val Leu Gln Glu Thr Leu Lys Ser Leu Gly Arg Tyr  
           35                  40          45  
 Arg Leu Leu Gly Lys Ile Phe Arg Lys Gln Glu Asn Ala Asn Ala Val  
       50                  55          60  
 Leu Leu Glu Leu Leu Glu Asp Thr Asp Val Ser Ala Ile Pro Ser Glu  
 65                  70          75          80  
 Val Gln Gly Lys Gly Gly Val Trp Lys Val Ile Phe Lys Thr Pro Asn  
                   85          90          95  
 Gln Asp Thr Glu Phe Leu Glu Arg Leu Asn Leu Phe Leu Glu Lys Glu  
           100                  105          110  
 Gly Gln Thr Val Ser Gly Met Phe Arg Ala Leu Gly Gln Glu Ala Leu  
           115                  120          125  
 Ser Pro Ala Thr Val Pro Cys Ile Ser Pro Glu Leu Leu Ala His Leu  
       130                  135          140  
 Leu Gly Gln Ala Met Ala His Ala Pro Gln Pro Leu Leu Pro Met Arg  
 145                  150          155          160  
 Tyr Arg Lys Leu Arg Val Phe Ser Gly Ser Ala Val Pro Ala Pro Glu  
                   165          170          175  
 Glu Glu Ser Phe Glu Val Trp Leu Glu Gln Ala Thr Glu Ile Val Lys  
           180                  185          190  
 Glu Trp Pro  
       195

<210> 8  
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 <212> DNA  
 <213> homo sapiens

<220>  
 <221> CDS  
 <222> (2)...(448)

<400> 8  
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   Asp Leu Met His Ile Val Gln Ala Asp Asn Pro Ser Ile Ser Val Glu  
    1                  5                  10                  15  
  
 gag tgt ttg gag gcc ttt aag caa gtg ttt ggg agc cta gag agc cgc 97  
 Glu Cys Leu Glu Ala Phe Lys Gln Val Phe Gly Ser Leu Glu Ser Arg  
           20                  25                  30  
  
 agg aca gcc cag gtg agg tat ctg aag ccc tat cag gag gaa gga gag 145  
 Arg Thr Ala Gln Val Arg Tyr Leu Lys Pro Tyr Gln Glu Glu Gly Glu  
           35                  40                  45  
  
 aag gtc tca gcc tat gtg tta cgg cta gaa acc ctg ctc cgg aga gcg 193  
 Lys Val Ser Ala Tyr Val Leu Arg Leu Glu Thr Leu Leu Arg Arg Ala  
           50                  55                  60  
  
 gtg gag aaa cgc gcc atc cct cgg cgt att gcg gac cag gtc cgc ctg 241  
 Val Glu Lys Arg Ala Ile Pro Arg Arg Ile Ala Asp Gln Val Arg Leu  
       65                  70                  75                  80  
  
 gag cag gtc atg gct ggg gcc act ctt aac cag atg ctg tgg tgc cgg 289  
 Glu Gln Val Met Ala Gly Ala Thr Leu Asn Gln Met Leu Trp Cys Arg  
           85                  90                  95  
  
 ctt agg gag ctg aag gat cag ggc ccg ccc ccc agc ttc ctt gag cta 337

Leu Arg Glu Leu Lys Asp Gln Gly Pro Pro Pro Ser Phe Leu Glu Leu  
 100 105 110  
 atg aag gta ata cgg gaa gaa gag gag gaa gag gcc tcc ttt gag aat 385  
 Met Lys Val Ile Arg Glu Glu Glu Glu Glu Glu Ala Ser Phe Glu Asn  
 115 120 125  
 gag agt atc gaa gag cca gag gaa cga gat ggc tat ggc cgc tgg aat 433  
 Glu Ser Ile Glu Glu Pro Glu Glu Arg Asp Gly Tyr Gly Arg Trp Asn  
 130 135 140  
 cat gag gga gac gac tgaaaaccac ctggggggcag gacccacagc cagtgggcta 488  
 His Glu Gly Asp Asp  
 145

agaccttttaa aaaatTTTTTT tctttaaTgt atgggactga aatcaaacca tgaaagccaa 548  
 ttattgacct tccttccttc ctctcttccc tcccttcctc cttctctcct tctctccttt 608  
 tttttttttt tttttaaac ctgttcttgg gtgggtgtgg gtataatact aagttgagat 668  
 gatattcattt acgggggaag gcgctttgtg aagtaggcct tatttctctt gtcctttcgt 728  
 acagggagga atttgaagta gatagaaacc gacctggatt actccggtct gaactcagat 788  
 cacgtaggac tttaatcggt gaacaaacga acctttaata gcggg 833

<210> 9  
 <211> 149  
 <212> PRT  
 <213> homo sapiens

<400> 9  
 Asp Leu Met His Ile Val Gln Ala Asp Asn Pro Ser Ile Ser Val Glu  
 1 5 10 15  
 Glu Cys Leu Glu Ala Phe Lys Gln Val Phe Gly Ser Leu Glu Ser Arg  
 20 25 30  
 Arg Thr Ala Gln Val Arg Tyr Leu Lys Pro Tyr Gln Glu Glu Gly Glu  
 35 40 45  
 Lys Val Ser Ala Tyr Val Leu Arg Leu Glu Thr Leu Leu Arg Arg Ala  
 50 55 60  
 Val Glu Lys Arg Ala Ile Pro Arg Arg Ile Ala Asp Gln Val Arg Leu  
 65 70 75 80  
 Glu Gln Val Met Ala Gly Ala Thr Leu Asn Gln Met Leu Trp Cys Arg  
 85 90 95  
 Leu Arg Glu Leu Lys Asp Gln Gly Pro Pro Pro Ser Phe Leu Glu Leu  
 100 105 110  
 Met Lys Val Ile Arg Glu Glu Glu Glu Glu Ala Ser Phe Glu Asn  
 115 120 125  
 Glu Ser Ile Glu Glu Pro Glu Glu Arg Asp Gly Tyr Gly Arg Trp Asn  
 130 135 140  
 His Glu Gly Asp Asp  
 145

<210> 10  
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 <212> DNA  
 <213> homo sapiens

<220>  
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 <222> (2)...(850)





|   |      |
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| ccc agc ttc ctt gag cta atg aag gta ata cgg gaa gaa gag gag gaa           | 769  |
| Pro Ser Phe Leu Glu Leu Met Lys Val Ile Arg Glu Glu Glu Glu Glu           |      |
| 245                                250                                255 |      |
| gag gcc tcc ttt gag aat gag agt atc gaa gag cca gag gaa cga gat           | 817  |
| Glu Ala Ser Phe Glu Asn Glu Ser Ile Glu Glu Pro Glu Glu Arg Asp           |      |
| 260                                265                                270 |      |
| ggc tat ggc cgc tgg aat cat gag gga gac gac tgaaaaccac ctggggggcgag       | 870  |
| Gly Tyr Gly Arg Trp Asn His Glu Gly Asp Asp                               |      |
| 275                                280                                    |      |
| gacctcacgc cagtgggcta agacctttaaa aaaattttttt tctttaaatgt atggggactga     | 930  |
| aatacaaacca tgaaagccaa ttatttgacct tccttccttc ctctccttcc ttcccttccct      | 990  |
| ccttctctcc ttctctcctc ctctctcctc tcctctcctc tctttccttc ctctccttcc         | 1050 |
| tctttctttt tctcttttct ctttatcttct tgggtctcac tctcatcacc caggctagag        | 1110 |
| tgcagtggca caaaaaatctc ggctcactgc agccttgact tcccaggctc aggctcaggt        | 1170 |
| gatcctcaca ccttagcctc ccaagtacct gggactacag gcacgcacca ccatgcctag         | 1230 |
| ctattctttt gtatttttgg tagagacagg gttttgctgt gttgctcagg ctgggtctgga        | 1290 |
| acccttaggc tcaaatgatg tgccaactc ggcctcccaa agtgctggga ttacaggcat          | 1350 |
| gaaccgccat gcctggccct tgatttttct ttttaagaaa aaaatatcta ggagtttctt         | 1410 |
| agaccctatg tagattatta atgaacaaaa gattaaactc caaatattaa atagtaagcc         | 1470 |
| tgaagggaatc tgaaacactt gtactttcaa ttttctttaa ataatcccaa atagaccaga        | 1530 |
| attggcccat accatagaag aaagaattgg cagtcaaaaa aaaa                          | 1574 |
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| <213> homo sapiens  |      |
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| Val Gln Gly Lys Gly Gly Val Trp Lys Val Ile Phe Lys Thr Pro Asn           |      |
| 1                    5                    10                    15        |      |
| Gln Asp Thr Glu Phe Leu Glu Arg Leu Asn Leu Phe Leu Glu Lys Glu           |      |
| 20                    25                    30                            |      |
| Gly Gln Thr Val Ser Gly Met Phe Arg Ala Leu Gly Gln Glu Gly Val           |      |
| 35                    40                    45                            |      |
| Ser Pro Ala Thr Val Pro Cys Ile Ser Pro Glu Leu Leu Ala His Leu           |      |
| 50                    55                    60                            |      |
| Leu Gly Gln Ala Met Ala His Ala Pro Gln Pro Leu Leu Pro Met Arg           |      |
| 65                    70                    75                    80      |      |
| Tyr Arg Lys Leu Arg Val Phe Ser Gly Ser Ala Val Pro Ala Pro Glu           |      |
| 85                    90                    95                            |      |
| Glu Glu Ser Phe Glu Val Trp Leu Glu Gln Ala Thr Glu Ile Val Lys           |      |
| 100                    105                    110                         |      |
| Glu Trp Pro Val Thr Glu Ala Glu Lys Lys Arg Trp Leu Ala Glu Ser           |      |
| 115                    120                    125                         |      |
| Leu Arg Gly Pro Ala Leu Asp Leu Met His Ile Val Gln Ala Asp Asn           |      |
| 130                    135                    140                         |      |
| Pro Ser Ile Ser Val Glu Glu Cys Leu Glu Ala Phe Lys Gln Val Phe           |      |
| 145                    150                    155                    160  |      |
| Gly Ser Leu Glu Ser Arg Arg Thr Ala Gln Val Arg Tyr Leu Lys Thr           |      |
| 165                    170                    175                         |      |
| Tyr Gln Glu Glu Gly Glu Lys Val Ser Ala Tyr Val Leu Arg Leu Glu           |      |
| 180                    185                    190                         |      |
| Thr Leu Leu Arg Lys Ala Val Glu Lys Arg Ala Ile Pro Arg Arg Ile           |      |
| 195                    200                    205                         |      |

Ala Asp Gln Val Arg Leu Glu Gln Val Met Ala Gly Ala Thr Leu Asn  
 210 215 220  
 Gln Met Leu Trp Cys Arg Leu Arg Glu Leu Lys Asp Gln Gly Pro Pro  
 225 230 235 240  
 Pro Ser Phe Leu Glu Leu Met Lys Val Ile Arg Glu Glu Glu Glu Glu  
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 Cys Arg Gly Glu His Leu Asn Thr Arg Arg Cys Met Leu Ile Leu Gly  
 10 15 20 25

atc ccc gag gac tgt ggc gag gat gag ttt gag gag aca ctc cag gag 150  
 Ile Pro Glu Asp Cys Gly Glu Asp Glu Phe Glu Glu Thr Leu Gln Glu  
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gct tgc agg cac ctg ggc aga tac agg gtg att ggc agg atg ttt agg 198  
 Ala Cys Arg His Leu Gly Arg Tyr Arg Val Ile Gly Arg Met Phe Arg  
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agg gag gag aac gcc cag gcg att cta ctg gag ctg gca caa gat atc 246  
 Arg Glu Glu Asn Ala Gln Ala Ile Leu Leu Glu Leu Ala Gln Asp Ile  
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gac tat gct ttg ctc cca agg gaa ata cca gga aag ggg ggg ccc tgg 294  
 Asp Tyr Ala Leu Leu Pro Arg Glu Ile Pro Gly Lys Gly Gly Pro Trp  
 75 80 85

gaa gtg att gta aaa ccc cgt aac tca gat ggg gaa ttt ctc aac aga 342  
 Glu Val Ile Val Lys Pro Arg Asn Ser Asp Gly Glu Phe Leu Asn Arg  
 90 95 100 105

ctg aac cgc ttc tta gag gag gag agg cgg acc gtg tca gat atg aac 390

|     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |     |      |  |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------|--|
| Leu | Asn | Arg | Phe | Leu | Glu | Glu | Glu | Arg | Arg | Thr | Val | Ser | Asp | Met | Asn |      |  |
|     |     |     |     | 110 |     |     |     |     | 115 |     |     |     |     | 120 |     |      |  |
| cga | gtc | ctc | ggg | tcg | gac | acc | aat | tgt | tcg | gct | cca | aga | gtg | act | ata | 438  |  |
| Arg | Val | Leu | Gly | Ser | Asp | Thr | Asn | Cys | Ser | Ala | Pro | Arg | Val | Thr | Ile |      |  |
|     |     |     | 125 |     |     |     |     | 130 |     |     |     |     | 135 |     |     |      |  |
| tca | cca | gag | ttc | tgg | acc | tgg | gcc | cag | act | ctg | ggg | gca | gca | gtg | cag | 486  |  |
| Ser | Pro | Glu | Phe | Trp | Thr | Trp | Ala | Gln | Thr | Leu | Gly | Ala | Ala | Val | Gln |      |  |
|     |     |     | 140 |     |     |     |     | 145 |     |     |     | 150 |     |     |     |      |  |
| cct | ctg | cta | gaa | caa | atg | ttg | tac | cga | gaa | cta | aga | gtg | ttt | tct | ggg | 534  |  |
| Pro | Leu | Leu | Glu | Gln | Met | Leu | Tyr | Arg | Glu | Leu | Arg | Val | Phe | Ser | Gly |      |  |
|     | 155 |     |     |     |     | 160 |     |     |     |     | 165 |     |     |     |     |      |  |
| aac | acc | ata | tcc | atc | cca | ggg | gca | ctg | gcc | ttt | gat | gcc | tgg | ctt | gag | 582  |  |
| Asn | Thr | Ile | Ser | Ile | Pro | Gly | Ala | Leu | Ala | Phe | Asp | Ala | Trp | Leu | Glu |      |  |
| 170 |     |     |     |     | 175 |     |     |     |     | 180 |     |     |     |     | 185 |      |  |
| cac | acc | act | gag | atg | cta | cag | atg | tgg | cag | gtg | ccc | gag | ggg | gaa | aag | 630  |  |
| His | Thr | Thr | Glu | Met | Leu | Gln | Met | Trp | Gln | Val | Pro | Glu | Gly | Glu | Lys |      |  |
|     |     |     |     | 190 |     |     |     |     | 195 |     |     |     |     | 200 |     |      |  |
| agg | cgg | agg | ctg | atg | gaa | tgc | tta | cgg | ggc | cct | gct | ctc | cag | gtg | gtc | 678  |  |
| Arg | Arg | Arg | Leu | Met | Glu | Cys | Leu | Arg | Gly | Pro | Ala | Leu | Gln | Val | Val |      |  |
|     |     |     | 205 |     |     |     |     | 210 |     |     |     |     | 215 |     |     |      |  |
| agt | ggg | ctc | cgg | gcc | agc | aat | gct | tcc | ata | act | gtg | gag | gag | tgc | ctg | 726  |  |
| Ser | Gly | Leu | Arg | Ala | Ser | Asn | Ala | Ser | Ile | Thr | Val | Glu | Glu | Cys | Leu |      |  |
|     |     | 220 |     |     |     |     | 225 |     |     |     |     | 230 |     |     |     |      |  |
| gct | gcc | ttg | cag | cag | gtg | ttc | gga | cct | gtg | gag | agc | cat | aaa | att | gcc | 774  |  |
| Ala | Ala | Leu | Gln | Gln | Val | Phe | Gly | Pro | Val | Glu | Ser | His | Lys | Ile | Ala |      |  |
|     |     | 235 |     |     |     | 240 |     |     |     |     | 245 |     |     |     |     |      |  |
| cag | gtg | aag | ttg | tgt | aaa | gcc | tat | cag | gag | gca | gga | gag | aaa | gta | tct | 822  |  |
| Gln | Val | Lys | Leu | Cys | Lys | Ala | Tyr | Gln | Glu | Ala | Gly | Glu | Lys | Val | Ser |      |  |
| 250 |     |     |     | 255 |     |     |     |     |     | 260 |     |     |     | 265 |     |      |  |
| agc | ttt | gtg | tta | cgt | ttg | gaa | ccc | ctg | ctc | caa | aga | gct | gta | gaa | aac | 870  |  |
| Ser | Phe | Val | Leu | Arg | Leu | Glu | Pro | Leu | Leu | Gln | Arg | Ala | Val | Glu | Asn |      |  |
|     |     |     |     | 270 |     |     |     |     | 275 |     |     |     |     | 280 |     |      |  |
| aat | gtg | gta | tca | cgt | aga | aac | gtg | aat | cag | act | cgc | ctg | aaa | cga | gtc | 918  |  |
| Asn | Val | Val | Ser | Arg | Arg | Asn | Val | Asn | Gln | Thr | Arg | Leu | Lys | Arg | Val |      |  |
|     |     |     | 285 |     |     |     |     | 290 |     |     |     |     | 295 |     |     |      |  |
| tta | agt | ggg | gcc | acc | ctt | cct | gac | aaa | ctc | cga | gat | aag | ctt | aag | ctg | 966  |  |
| Leu | Ser | Gly | Ala | Thr | Leu | Pro | Asp | Lys | Leu | Arg | Asp | Lys | Leu | Lys | Leu |      |  |
|     |     | 300 |     |     |     |     | 305 |     |     |     |     | 310 |     |     |     |      |  |
| atg | aaa | cag | cga | agg | aag | cct | cct | ggg | ttc | ctg | gcc | ctg | gtg | aag | ctc | 1014 |  |
| Met | Lys | Gln | Arg | Arg | Lys | Pro | Pro | Gly | Phe | Leu | Ala | Leu | Val | Lys | Leu |      |  |
|     |     | 315 |     |     |     | 320 |     |     |     |     | 325 |     |     |     |     |      |  |
| ctg | cgt | gag | gag | gag | gaa | tgg | gag | gcc | act | tta | ggg | cca | gat | agg | gag | 1062 |  |
| Leu | Arg | Glu | Glu | Glu | Glu | Trp | Glu | Ala | Thr | Leu | Gly | Pro | Asp | Arg | Glu |      |  |
| 330 |     |     |     |     | 335 |     |     |     |     | 340 |     |     |     |     | 345 |      |  |

agt ctg gag ggg ctg gaa gta gcc cca agg cca cct gcc agg atc act 1110  
 Ser Leu Glu Gly Leu Glu Val Ala Pro Arg Pro Pro Ala Arg Ile Thr  
 350 355 360

ggg gtt ggg gca gta cct ctc cct gcc tct ggc aac agt ttt gat gcg 1158  
 Gly Val Gly Ala Val Pro Leu Pro Ala Ser Gly Asn Ser Phe Asp Ala  
 365 370 375

agg cct tcc cag ggc tac cgg cgc cgg agg ggc aga ggc caa cac cga 1206  
 Arg Pro Ser Gln Gly Tyr Arg Arg Arg Gly Arg Gly Gln His Arg  
 380 385 390

agg ggt ggt gtg gca agg gct ggc tct cga ggc tca aga aaa cgg aaa 1254  
 Arg Gly Gly Val Ala Arg Ala Gly Ser Arg Gly Ser Arg Lys Arg Lys  
 395 400 405

cgc cac aca ttc tgc tat agc tgt ggg gaa gac ggc cac atc agg gta 1302  
 Arg His Thr Phe Cys Tyr Ser Cys Gly Glu Asp Gly His Ile Arg Val  
 410 415 420 425

cag tgc atc aac ccc tcc aac ctg ctc ttg gta aag cag aag aaa cag 1350  
 Gln Cys Ile Asn Pro Ser Asn Leu Leu Leu Val Lys Gln Lys Lys Gln  
 430 435 440

gct gca gtt gag tcg gga aac ggg aac tgg gct tgg gac aag agc cat 1398  
 Ala Ala Val Glu Ser Gly Asn Gly Asn Trp Ala Trp Asp Lys Ser His  
 445 450 455

ccc aag tcc aag gcc aag taggctcggg agaacagggc aacatttcct 1446  
 Pro Lys Ser Lys Ala Lys  
 460

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 Asp Glu Phe Glu Glu Thr Leu Gln Glu Ala Cys Arg His Leu Gly Arg  
 35 40 45

Tyr Arg Val Ile Gly Arg Met Phe Arg Arg Glu Glu Asn Ala Gln Ala  
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 Ile Leu Leu Glu Leu Ala Gln Asp Ile Asp Tyr Ala Leu Leu Pro Arg  
 65 70 75 80  
 Glu Ile Pro Gly Lys Gly Gly Pro Trp Glu Val Ile Val Lys Pro Arg  
 85 90 95  
 Asn Ser Asp Gly Glu Phe Leu Asn Arg Leu Asn Arg Phe Leu Glu Glu  
 100 105 110  
 Glu Arg Arg Thr Val Ser Asp Met Asn Arg Val Leu Gly Ser Asp Thr  
 115 120 125  
 Asn Cys Ser Ala Pro Arg Val Thr Ile Ser Pro Glu Phe Trp Thr Trp  
 130 135 140  
 Ala Gln Thr Leu Gly Ala Ala Val Gln Pro Leu Leu Glu Gln Met Leu  
 145 150 155 160  
 Tyr Arg Glu Leu Arg Val Phe Ser Gly Asn Thr Ile Ser Ile Pro Gly  
 165 170 175  
 Ala Leu Ala Phe Asp Ala Trp Leu Glu His Thr Thr Glu Met Leu Gln  
 180 185 190  
 Met Trp Gln Val Pro Glu Gly Glu Lys Arg Arg Arg Leu Met Glu Cys  
 195 200 205  
 Leu Arg Gly Pro Ala Leu Gln Val Val Ser Gly Leu Arg Ala Ser Asn  
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 Ala Ser Ile Thr Val Glu Glu Cys Leu Ala Ala Leu Gln Gln Val Phe  
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 Gly Pro Val Glu Ser His Lys Ile Ala Gln Val Lys Leu Cys Lys Ala  
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 Tyr Gln Glu Ala Gly Glu Lys Val Ser Ser Phe Val Leu Arg Leu Glu  
 260 265 270  
 Pro Leu Leu Gln Arg Ala Val Glu Asn Asn Val Val Ser Arg Arg Asn  
 275 280 285  
 Val Asn Gln Thr Arg Leu Lys Arg Val Leu Ser Gly Ala Thr Leu Pro  
 290 295 300  
 Asp Lys Leu Arg Asp Lys Leu Lys Leu Met Lys Gln Arg Arg Lys Pro  
 305 310 315 320  
 Pro Gly Phe Leu Ala Leu Val Lys Leu Leu Arg Glu Glu Glu Glu Trp  
 325 330 335  
 Glu Ala Thr Leu Gly Pro Asp Arg Glu Ser Leu Glu Gly Leu Glu Val  
 340 345 350  
 Ala Pro Arg Pro Pro Ala Arg Ile Thr Gly Val Gly Ala Val Pro Leu  
 355 360 365  
 Pro Ala Ser Gly Asn Ser Phe Asp Ala Arg Pro Ser Gln Gly Tyr Arg  
 370 375 380  
 Arg Arg Arg Gly Arg Gly Gln His Arg Arg Gly Gly Val Ala Arg Ala  
 385 390 395 400  
 Gly Ser Arg Gly Ser Arg Lys Arg Lys Arg His Thr Phe Cys Tyr Ser  
 405 410 415  
 Cys Gly Glu Asp Gly His Ile Arg Val Gln Cys Ile Asn Pro Ser Asn  
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| cccgcaaagg | ctcccgacc  | tctgcgtgtt | aaagagacga | gcacgcacat | cactgtaagc | 120 |
| ggcggcgcgg | cggcgggcct | ggtcgaatta | gaatttaa   | actctgagca | ccatgacact | 180 |
| gagacttcta | gaagactggt | gcagagggat | ggatatgaat | cctcggaaag | cactattggt | 240 |
| tgccggcatc | cctccgacct | gcggagtggc | agacatagag | gaggccctgc | aggctggcct | 300 |
| tgctccctta | ggggaacaca | gactgcttgg | gaggatgttc | aggagggatg | agaacaagaa | 360 |
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| gcaaaaggag | gtgtctggag | agtgatcttt | aagcctcctg | atactgatag | tgactttttg | 480 |
| tcagattaa  | atgagttttt | aaagggggag | ggcatgacga | tgggtgaatt |            | 530 |

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